Serial Number: ___09/8○○○089 B___

Processing Date: 5-19-03

Edited by: _M. Scholar_

Verified by: _____ (STIC staff) #16

ENTERED

☐ Changed a file from non-ASCII to ASCII

☑ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☑ Edited a format error in the Current Application Data section, specifically _____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: _____

☐ Deleted extra, invalid, headings used by an applicant, specifically: _____

☑ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically: _____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: _____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other: _____

_____

_____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

RAW SEQUENCE LISTING   DATE: 05/19/2003
PATENT APPLICATION: **US/09/870,089B**   TIME: 14:15:29

Input Set : **A:\ptoms.txt**
Output Set: **N:\CRF4\05192003\I870089B.raw**

```
 3 <110> APPLICANT: Charles A. Nicolette
 5 <120> TITLE OF INVENTION: THERAPEUTIC COMPOUNDS FOR OVARIAN CANCER
 8 <130> FILE REFERENCE: 68126881209900
10 <140> CURRENT APPLICATION NUMBER: 09/870,089B
11 <141> CURRENT FILING DATE: 2001-05-30
13 <160> NUMBER OF SEQ ID NOS: 14
15 <170> SOFTWARE: FastSEQ for Windows Version 3.0
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 2015
19 <212> TYPE: DNA
20 <213> ORGANISM: Homo sapiens
22 <400> SEQUENCE: 1
23  gttttctact ttgcccgccc acagatgtag ttttctctgc gcgtgtgcgt tttccctcct     60
24  cccccgccct cagggtccac ggccaccatg gcgtattagg ggcagcagtg cctgcggcag    120
25  cattggcctt tgcagcggcg gcagcagcac caggctctgc agcggcaacc cccagcggct    180
26  taagccatgg cgtgagtacc ggggcgggtc gtccagctgt gctcctgggg ccggcgcggg    240
27  ttttggattg gtgggggtgcg gcctggggcc agggcggtgc cgccaagggg gaagcgattt    300
28  aacgagcgcc cgggacgcgt ggtctttgct tgggtgtccc cgagacgctc gcgtgcctgg    360
29  gatcgggaaa gcgtagtcgg gtgcccggac tgcttcccca ggagccctac agccctcgga    420
30  ccccgagccc cgcaaggtcc caggggtctt ggctgttgcc ccacgaaacg tgcaggaacc    480
31  aagatggcgg cggcagggcg gcggcgcggg cgtgagtcaa gggcgggcgg tgggcggggc    540
32  gcggccgctg gccgtatttg gacgtgggga cggagcgctt tcctcttggc ggccggtgga    600
33  agaatcccct ggtctccgtg agcgtccatt ttgtggaacc tgagttgcaa gcagggaggg    660
34  gcaaatacaa ctgccctgtt cccgattctc tagatggccg atctagagaa gtcccgcctc    720
35  ataagtggaa ggatgaaatt ctcagaacag ctaacctcta atgggagttg gcttctgatt    780
36  ctcattcagg cttctcacgg cattcagcag cagcgttgct gtaaccgaca aagacacctt    840
37  cgaattaagc acattcctcg attccagcaa agcaccgcaa catgaccgaa atgagcttcc    900
38  tgagcagcga ggtgttggtg ggggacttga tgtccccctt cgaccgtcg ggtttggggg     960
39  ctgaagaaag cctaggtctc ttagatgatt acctggaggt ggccaagcac ttcaaacctc   1020
40  atgggttctc cagcgacaag gctaaggcgg gctcctccga atggctggct gtggatgggt   1080
41  tggtcagtcc ctccaacaac agcaaggagg atgccttctc cgggacagat tggatgttgg   1140
42  agaaaatgga tttgaaggag ttcgacttgg atgccctgtt gggtatagat gacctggaaa   1200
43  ccatgccaga tgaccttctg accacgttgg atgacacttg tgatctcttt gcccccctag   1260
44  tccaggagac taataagcag cccccccaga cggtgaaccc aattggccat ctcccagaaa   1320
45  gtttaacaaa acccgaccag gttgccccct tcaccttctt acaacctctt cccctttccc   1380
46  caggggtcct gtcctccact ccagatcatt cctttagttt agagctgggc agtgaagtgg   1440
47  atatcactga aggagatagg aagccagact acactgctta cgttgccatg atccctcagt   1500
48  gcataaagga ggaagacacc ccttcagata atgatagtgg catctgtatg agcccagagt   1560
49  cctatctggg gtctcctcag cacagcccct ctaccagggg ctctccaaat aggagcctcc   1620
50  catctccagg tgttctctgt gggtctgccc gtcccaaacc ttacgatcct cctggagaga   1680
51  agatggtagc agcaaaagta aagggtgaga aactggataa gaagctgaaa aaaatggagc   1740
52  aaaacaagac agcagccact aggtaccgcc agaagaagag ggcggagcag gaggctctta   1800
53  ctggtgagtg caaagagctg gaaaagaaga cgaggctct aaaagagagg gcggattccc   1860
```

RAW SEQUENCE LISTING   DATE: 05/19/2003
PATENT APPLICATION: **US/09/870,089B**   TIME: 14:15:29

Input Set : **A:\ptoms.txt**
Output Set: **N:\CRF4\05192003\I870089B.raw**

```
54  tggccaagga gatccagtac ctgaaagatt tgatagaaga ggtccgcaag gcaaggggga   1920
55  agaaaagggt cccctagttg aggatagtca ggagcgtcaa tgtgcttgta catagagtgc   1980
56  tgtagctgtg tgttccaata aattattttg taggg                             2015
58 <210> SEQ ID NO: 2
59 <211> LENGTH: 351
60 <212> TYPE: PRT
61 <213> ORGANISM: Homo sapiens
63 <400> SEQUENCE: 2
64  Met Thr Glu Met Ser Phe Leu Ser Ser Glu Val Leu Val Gly Asp Leu
65  1               5                   10                  15
66  Met Ser Pro Phe Asp Pro Ser Gly Leu Gly Ala Glu Glu Ser Leu Gly
67              20                  25                  30
68  Leu Leu Asp Asp Tyr Leu Glu Val Ala Lys His Phe Lys Pro His Gly
69          35                  40                  45
70  Phe Ser Ser Asp Lys Ala Lys Ala Gly Ser Ser Glu Trp Leu Ala Val
71      50                  55                  60
72  Asp Gly Leu Val Ser Pro Ser Asn Asn Ser Lys Glu Asp Ala Phe Ser
73  65                  70                  75                  80
74  Gly Thr Asp Trp Met Leu Glu Lys Met Asp Leu Lys Glu Phe Asp Leu
75              85                  90                  95
76  Asp Ala Leu Leu Gly Ile Asp Asp Leu Glu Thr Met Pro Asp Asp Leu
77          100                 105                 110
78  Leu Thr Thr Leu Asp Asp Thr Cys Asp Leu Phe Ala Pro Leu Val Gln
79          115                 120                 125
80  Glu Thr Asn Lys Gln Pro Pro Gln Thr Val Asn Pro Ile Gly His Leu
81      130                 135                 140
82  Pro Glu Ser Leu Thr Lys Pro Asp Gln Val Ala Pro Phe Thr Phe Leu
83  145                 150                 155                 160
84  Gln Pro Leu Pro Leu Ser Pro Gly Val Leu Ser Ser Thr Pro Asp His
85              165                 170                 175
86  Ser Phe Ser Leu Glu Leu Gly Ser Glu Val Asp Ile Thr Glu Gly Asp
87              180                 185                 190
88  Arg Lys Pro Asp Tyr Thr Ala Tyr Val Ala Met Ile Pro Gln Cys Ile
89          195                 200                 205
90  Lys Glu Glu Asp Thr Pro Ser Asp Asn Asp Ser Gly Ile Cys Met Ser
91      210                 215                 220
92  Pro Glu Ser Tyr Leu Gly Ser Pro Gln His Ser Pro Ser Thr Arg Gly
93  225                 230                 235                 240
94  Ser Pro Asn Arg Ser Leu Pro Ser Pro Gly Val Leu Cys Gly Ser Ala
95              245                 250                 255
96  Arg Pro Lys Pro Tyr Asp Pro Pro Gly Glu Lys Met Val Ala Ala Lys
97              260                 265                 270
98  Val Lys Gly Glu Lys Leu Asp Lys Lys Leu Lys Lys Met Glu Gln Asn
99          275                 280                 285
100  Lys Thr Ala Ala Thr Arg Tyr Arg Gln Lys Lys Arg Ala Glu Gln Glu
101          290                 295                 300
102  Ala Leu Thr Gly Glu Cys Lys Glu Leu Glu Lys Lys Asn Glu Ala Leu
103  305                 310                 315                 320
104  Lys Glu Arg Ala Asp Ser Leu Ala Lys Glu Ile Gln Tyr Leu Lys Asp
```

```
105                   325            330                 335
106   Leu Ile Glu Glu Val Arg Lys Ala Arg Gly Lys Lys Arg Val Pro
107             340              345              350
109 <210> SEQ ID NO: 3
110 <211> LENGTH: 9
111 <212> TYPE: PRT
112 <213> ORGANISM: Artificial Sequence
114 <220> FEATURE:
115 <223> OTHER INFORMATION: ATF4/CREB-2
117 <400> SEQUENCE: 3
118   Phe Leu Tyr Lys Trp His Gly Phe Val
119   1               5
121 <210> SEQ ID NO: 4
122 <211> LENGTH: 27
123 <212> TYPE: DNA
124 <213> ORGANISM: Artificial Sequence
126 <220> FEATURE:
127 <223> OTHER INFORMATION: ATF4/CREB-2
```
W--> `129 <221> NAME/KEY: misc_feature`
```
130 <222> LOCATION: (6)...(6)
131 <223> OTHER INFORMATION: n is A, C, G or T
```
W--> `133 <221> misc_feature`
```
134 <222> LOCATION: (21)...(21)
135 <223> OTHER INFORMATION: n is A, C, G or T
```
W--> `137 <221> misc_feature`
```
138 <222> LOCATION: (27)...(27)
139 <223> OTHER INFORMATION: n is A, C, G or T
```
W--> `141 <400> 4`
W--> `142   ttyytntaya artggcaygg nttygtn                                    27`
```
144 <210> SEQ ID NO: 5
145 <211> LENGTH: 9
146 <212> TYPE: PRT
147 <213> ORGANISM: Artificial Sequence
149 <220> FEATURE:
150 <223> OTHER INFORMATION: ATF4/CREB-2
152 <400> SEQUENCE: 5
153   Phe Leu His Lys Val His Phe Tyr Val
154   1               5
156 <210> SEQ ID NO: 6
157 <211> LENGTH: 27
158 <212> TYPE: DNA
159 <213> ORGANISM: Artificial Sequence
161 <220> FEATURE:
162 <223> OTHER INFORMATION: ATF4/CREB-2
```
W--> `164 <221> NAME/KEY: misc_feature`
```
165 <222> LOCATION: (6)...(6)
166 <223> OTHER INFORMATION: n is A, C, G or T
```
W--> `168 <221> misc_feature`
```
169 <222> LOCATION: (15)...(15)
```

```
       170 <223> OTHER INFORMATION: n is A, C, G or T
W-->   172 <221> misc_feature
       173 <222> LOCATION: (27)...(27)
       174 <223> OTHER INFORMATION: n is A, C, G or T
W-->   176 <400> 6
W-->   177  ttyytncaya argtncaytt ytaygtn                              27
       179 <210> SEQ ID NO: 7
       180 <211> LENGTH: 9
       181 <212> TYPE: PRT
       182 <213> ORGANISM: Artificial Sequence
       184 <220> FEATURE:
       185 <223> OTHER INFORMATION: ATF4/CREB-2
       187 <400> SEQUENCE: 7
       188  Phe Leu His Lys Trp His Trp Val Val
       189    1                5
       191 <210> SEQ ID NO: 8
       192 <211> LENGTH: 27
       193 <212> TYPE: DNA
       194 <213> ORGANISM: Artificial Sequence
       196 <220> FEATURE:
       197 <223> OTHER INFORMATION: ATF4/CREB-2
W-->   199 <221> NAME/KEY: misc_feature
       200 <222> LOCATION: (6)...(6)
       201 <223> OTHER INFORMATION: n is A, C, G or T
W-->   203 <221> misc_feature
       204 <222> LOCATION: (24)...(24)
       205 <223> OTHER INFORMATION: n is A, C, G or T
W-->   207 <221> misc_feature
       208 <222> LOCATION: (27)...(27)
       209 <223> OTHER INFORMATION: n is A, C, G or T
W-->   211 <400> 8
W-->   212  ttyytncaya artggcaytg ggtngtn                              27
       214 <210> SEQ ID NO: 9
       215 <211> LENGTH: 9
       216 <212> TYPE: PRT
       217 <213> ORGANISM: Artificial Sequence
       219 <220> FEATURE:
       220 <223> OTHER INFORMATION: ATF4/CREB-2
       222 <400> SEQUENCE: 9
       223  Phe Leu His Lys Trp His Trp Tyr Val
       224    1                5
       226 <210> SEQ ID NO: 10
       227 <211> LENGTH: 27
       228 <212> TYPE: DNA
       229 <213> ORGANISM: Artificial Sequence
       231 <220> FEATURE:
       232 <223> OTHER INFORMATION: ATF4/CREB-2
W-->   234 <221> NAME/KEY: misc_feature
       235 <222> LOCATION: (6)...(6)
```

```
       236 <223> OTHER INFORMATION: n is A, C, G or T
W--> 238 <221> misc_feature
       239 <222> LOCATION: (27)...(27)
       240 <223> OTHER INFORMATION: n is A, C, G or T
W--> 242 <400> 10
W--> 243    ttyytncaya artggcaytg gtaygtn                                    27
       245 <210> SEQ ID NO: 11
       246 <211> LENGTH: 9
       247 <212> TYPE: PRT
       248 <213> ORGANISM: Artificial Sequence
       250 <220> FEATURE:
       251 <223> OTHER INFORMATION: ATF4/CREB-2
       253 <400> SEQUENCE: 11
       254   Phe Leu His Lys Val His Tyr Leu Val
       255    1                 5
       257 <210> SEQ ID NO: 12
       258 <211> LENGTH: 27
       259 <212> TYPE: DNA
       260 <213> ORGANISM: Artificial Sequence
       262 <220> FEATURE:
       263 <223> OTHER INFORMATION: ATF4/CREB-2
W--> 265 <221> NAME/KEY: misc_feature
       266 <222> LOCATION: (6)...(6)
       267 <223> OTHER INFORMATION: n is A, C, G or T
W--> 269 <221> misc_feature
       270 <222> LOCATION: (15)...(15)
       271 <223> OTHER INFORMATION: n is A, C, G or T
W--> 273 <221> misc_feature
       274 <222> LOCATION: (22)...(22)
       275 <223> OTHER INFORMATION: n is A, C, G or T
W--> 277 <221> misc_feature
       278 <222> LOCATION: (24)...(24)
       279 <223> OTHER INFORMATION: n is A, C, G or T
W--> 281 <221> misc_feature
       282 <222> LOCATION: (27)...(27)
       283 <223> OTHER INFORMATION: n is A, C, G or T
W--> 285 <400> 12
W--> 286    ttyytncaya argtncayta yntngtn                                    27
       288 <210> SEQ ID NO: 13
       289 <211> LENGTH: 9
       290 <212> TYPE: PRT
       291 <213> ORGANISM: Homo sapiens
       293 <220> FEATURE:
       294 <223> OTHER INFORMATION: ATF4/CREB-2
       296 <400> SEQUENCE: 13
       297   Lys His Phe Lys Pro His Gly Phe Ser
       298    1                 5
       301 <210> SEQ ID NO: 14
       302 <211> LENGTH: 27  ·
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 05/19/2003
PATENT APPLICATION: US/09/870,089B   TIME: 14:15:30

Input Set : **A:\ptoms.txt**
Output Set: **N:\CRF4\05192003\I870089B.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; N Pos. 6,21,27
Seq#:6; N Pos. 6,15,27
Seq#:8; N Pos. 6,24,27
Seq#:10; N Pos. 6,27
Seq#:12; N Pos. 6,15,22,24,27
Seq#:14; N Pos. 15,21,27

VERIFICATION SUMMARY   DATE: 05/19/2003
PATENT APPLICATION: **US/09/870,089B**   TIME: 14:15:30

Input Set : **A:\ptoms.txt**
Output Set: **N:\CRF4\05192003\I870089B.raw**

```
L:129 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:133 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:137 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:141 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:142 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:164 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:168 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:172 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:176 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:199 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:203 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:207 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:211 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:234 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:238 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:242 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:265 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:269 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:273 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:277 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:281 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:285 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:309 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:313 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:317 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:321 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:322 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
```

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: **US/09/870,089B**

DATE: 05/07/2003
TIME: 11:01:17

Input Set : **A:\2099.00 Seq..txt**
Output Set: **N:\CRF4\05072003\I870089B.raw**

```
 3 <110> APPLICANT: Charles A. Nicolette
 5 <120> TITLE OF INVENTION: THERAPEUTIC COMPOUNDS FOR OVARIAN CANCER
 8 <130> FILE REFERENCE: 68126881209900
10 <140> CURRENT APPLICATION NUMBER: 09/870,089B
11 <141> CURRENT FILING DATE: 2001-05-30
13 <160> NUMBER OF SEQ ID NOS: 14
15 <170> SOFTWARE: FastSEQ for Windows Version 3.0
```

ERRORED SEQUENCES

```
        301 <210> SEQ ID NO: 14
        302 <211> LENGTH: 27
        303 <212> TYPE: DNA
        304 <213> ORGANISM: Artificial Sequence
        306 <220> FEATURE:
        307 <223> OTHER INFORMATION: ATF4/CREB-2
W--> 309 <221> NAME/KEY: misc_feature
        310 <222> LOCATION: (15)...(15)
        311 <223> OTHER INFORMATION: n is A, C, G or T
W--> 313 <221> misc_feature
        314 <222> LOCATION: (21)...(21)
        315 <223> OTHER INFORMATION: n is A, C, G or T
W--> 317 <221> misc_feature
        318 <222> LOCATION: (27)...(27)
        319 <223> OTHER INFORMATION: n is A, C, G or T
W--> 321 <400> 14
W--> 322  aarcayttya arccncaygg nttywsn                               27
E--> 325 r
```

remove extra material at end of file

VERIFICATION SUMMARY DATE: 05/07/2003
PATENT APPLICATION: **US/09/870,089B** TIME: 11:01:18

Input Set : **A:\2099.00 Seq..txt**
Output Set: **N:\CRF4\05072003\I870089B.raw**

```
L:129 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:133 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:137 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:141 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:142 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:164 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:168 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:172 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:176 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:199 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:203 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:207 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:211 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:234 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:238 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:242 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:265 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:269 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:273 M:258 W: Mandatory Feature missing, <220> Tag not· found for SEQ ID#:12
L:277 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:281 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:285 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:309 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:313 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:317 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:321 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:322 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
L:325 M:254 E: No. of Bases conflict, this line has no nucleotides.
```